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<110> Ross, Theodora
Mizukami, Ikuko

<120> Humoral Response to H1P1 in Cancer

<130> UM-08737

<140> 10/767,325

<141> 2004-01-29

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Leu Gly Thr His His Glu Lys Gly Ala Gln Thr Phe Trp Ser Val Val
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Asn Arg Leu Pro Leu Ser Ser Asn Ala Val Leu Cys Trp Lys Phe Cys
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His Val Phe His Lys Leu Leu Arg Asp Gly His Pro Asn Val Leu Lys
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Gly Tyr Leu Ser Glu Gly Tyr Gly Gln Leu Cys Ser Ile Tyr Leu Lys
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Gly Asn Leu Gln Met Ser Asp Arg Gln Leu Asp Glu Ala Gly Glu Ser
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Leu Ile Gln Val Ile Leu Asp Cys Ser His Leu Tyr Asp Tyr Thr Val
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Lys Leu Leu Phe Lys Leu His Ser Cys Leu Pro Ala Asp Thr Leu Gln
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Gly His Arg Asp Arg Phe Met Glu Gln Phe Thr Lys Leu Lys Asp Leu
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Phe Tyr Arg Ser Ser Asn Leu Gln Tyr Phe Lys Arg Leu Ile Gln Ile
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cccatatgtc agtgtaaate cttgttacct atcbcggtgtg tggtatttcc ccagccacag	2640
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Cys Arg Leu Ala Pro Leu Ile Gln Val Ile Leu Asp Cys Ser His Leu
 35 40 45

Tyr Asp Tyr Thr Val Lys Leu Leu Phe Lys Leu His Ser Cys Leu Pro
 50 55 60

Ala Asp Thr Leu Gln Gly His Arg Asp Arg Phe Met Glu Gln Phe Thr
 65 70 75 80

Lys Leu Lys Asp Leu Phe Tyr Arg Ser Ser Asn Leu Gln Tyr Phe Lys
 85 90 95

Arg Leu Ile Gln Ile Pro Gln Leu Pro Glu Asn Pro Pro Asn Phe Leu
 100 105 110

Arg Ala Ser Ala Leu Ser Glu His Ile Ser Pro Val Val Val Ile Pro
 115 120 125

Ala Glu Ala Ser Ser Pro Asp Ser Glu Pro Val Leu Glu Lys Asp Asp
 130 135 140

Leu Met Asp Met Asp Ala Ser Gln Gln Asn Leu Phe Asp Asn Lys Phe
 145 150 155 160

Asp Asp Ile Phe Gly Ser Ser Phe Ser Ser Asp Pro Phe Asn Phe Asn
 165 170 175

Ser Gln Asn Gly Val Asn Lys Asp Glu Lys Asp His Leu Ile Glu Arg
 180 185 190

Leu Tyr Arg Glu Ile Ser Gly Leu Lys Ala Gln Leu Glu Asn Met Lys
 195 200 205

Thr Glu Ser Gln Arg Val Val Leu Gln Leu Lys Gly His Val Ser Glu
 210 215 220

Leu Glu Ala Asp Leu Ala Glu Gln Gln His Leu Arg Gln Gln Ala Ala
 225 230 235 240

Asp Asp Cys Glu Phe Leu Arg Ala Glu Leu Asp Glu Leu Arg Arg Gln
 245 250 255

Arg Glu Asp Thr Glu Lys Ala Gln Arg Ser Leu Ser Glu Ile Glu Arg
 260 265 270

Lys Ala Gln Ala Asn Glu Gln Arg Tyr Ser Lys Leu Lys Glu Lys Tyr
 275 280 285

Ser Glu Leu Val Gln Asn His Ala Asp Leu Leu Arg Lys Asn Ala Glu
 290 295 300

Val Thr Lys Gln Val Ser Met Ala Arg Gln Ala Gln Val Asp Leu Glu
 305 310 315 320

Arg Glu Lys Lys Glu Leu Glu Asp Ser Leu Glu Arg Ile Ser Asp Gln
 325 330 335

Gly Gln Arg Lys Thr Gln Glu Gln Leu Glu Val Leu Glu Ser Leu Lys
 340 345 350

Gln Glu Leu Ala Thr Ser Gln Arg Glu Leu Gln Val Leu Gln Gly Ser
 355 360 365

Leu Glu Thr Ser Ala Gln Ser Glu Ala Asn Trp Ala Ala Glu Phe Ala
 370 375 380

Glu Leu Glu Lys Glu Arg Asp Ser Leu Val Ser Gly Ala Ala His Arg
 385 390 395 400

Glu Glu Glu Leu Ser Ala Leu Arg Lys Glu Leu Gln Asp Thr Gln Leu
 405 410 415

Lys Leu Ala Ser Thr Glu Glu Ser Met Cys Gln Leu Ala Lys Asp Gln
 420 425 430

Arg Lys Met Leu Leu Val Gly Ser Arg Lys Ala Ala Glu Gln Val Ile
 435 440 445

Gln Asp Ala Leu Asn Gln Leu Glu Glu Pro Pro Leu Ile Ser Cys Ala
 450 455 460

Gly Ser Ala Asp His Leu Leu Ser Thr Val Thr Ser Ile Ser Ser Cys
 465 470 475 480

Ile Glu Gln Leu Glu Lys Ser Trp Ser Gln Tyr Leu Ala Cys Pro Glu
 485 490 495

Asp Ile Ser Gly Leu Leu His Ser Ile Thr Leu Leu Ala His Leu Thr
 500 505 510

Ser Asp Ala Ile Ala His Gly Ala Thr Thr Cys Leu Arg Ala Pro Pro
 515 520 525

Glu Pro Ala Asp Ser Leu Thr Glu Ala Cys Lys Gln Tyr Gly Arg Glu
 530 535 540

Thr Leu Ala Tyr Leu Ala Ser Leu Glu Glu Glu Gly Ser Leu Glu Asn
 545 550 555 560

Ala Asp Ser Thr Ala Met Arg Asn Cys Leu Ser Lys Ile Lys Ala Ile
 565 570 575

Gly Glu Glu Leu Leu Pro Arg Gly Leu Asp Ile Lys Gln Glu Glu Leu
 580 585 590

Gly Asp Leu Val Asp Lys Glu Met Ala Ala Thr Ser Ala Ala Ile Glu
 595 600 605

Thr Ala Thr Ala Arg Ile Glu Glu Met Leu Ser Lys Ser Arg Ala Gly
 610 615 620

Asp Thr Gly Val Lys Leu Glu Val Asn Glu Arg Ile Leu Gly Cys Cys
 625 630 635 640

Thr Ser Leu Met Gln Ala Ile Gln Val Leu Ile Val Ala Ser Lys Asp
 645 650 655

Leu Gln Arg Glu Ile Val Glu Ser Gly Arg Gly Thr Ala Ser Pro Lys
 660 665 670

Glu Phe Tyr Ala Lys Asn Ser Arg Trp Thr Glu Gly Leu Ile Ser Ala
 675 680 685
 Ser Lys Ala Val Gly Trp Gly Ala Thr Val Met Val Asp Ala Ala Asp
 690 695 700
 Leu Val Val Gln Gly Arg Gly Lys Phe Glu Glu Leu Met Val Cys Ser
 705 710 715 720
 His Glu Ile Ala Ala Ser Thr Ala Gln Leu Val Ala Ala Ser Lys Val
 725 730 735
 Lys Ala Asp Lys Asp Ser Pro Asn Leu Ala Gln Leu Gln Gln Ala Ser
 740 745 750
 Arg Gly Val Asn Gln Ala Thr Ala Gly Val Val Ala Ser Thr Ile Ser
 755 760 765
 Gly Lys Ser Gln Ile Glu Glu Thr Asp Asn Met Asp Phe Ser Ser Met
 770 775 780
 Thr Leu Thr Gln Ile Lys Arg Gln Glu Met Asp Ser Gln Val Arg Val
 785 790 795 800
 Leu Glu Leu Glu Asn Glu Leu Gln Lys Glu Arg Gln Lys Leu Gly Glu
 805 810 815
 Leu Arg Lys Lys His Tyr Glu Leu Ala Gly Val Ala Glu Gly Trp Glu
 820 825 830
 Glu Gly Thr Glu Ala Ser Pro Pro Thr Leu Gln Glu Val Val Thr Glu
 835 840 845
 Lys Glu Ser Gln Thr Asn Thr Pro Tyr Val Ser Val Asn Pro Cys Tyr
 850 855 860
 Leu Ser Arg Val Cys Tyr Phe Pro Ser His Arg Pro Asn Pro Trp Ser
 865 870 875 880
 Pro Arg Gly Ser His Thr Thr Ala Ile Thr Gln Cys Arg Gly His Ala
 885 890 895
 His Phe Gln Arg Leu Pro Pro Arg His Pro Phe Cys Leu Asp Pro Trp
 900 905 910

Ile Ser Thr Ala Ser Tyr Gly Gly Trp Leu Gly Phe Leu Val Leu Phe
 915 920 925

Phe Phe Phe Lys Phe His Ser His Ser Gln Leu Ser Gln Arg Ala His
 930 935 940

Pro Trp Gly Val Ser Arg Ala Pro Gln Leu Trp Leu Gln Arg Trp Cys
 945 950 955 960

Cys Pro Gly Leu Ser Val Leu His Leu Arg Leu His Thr Asp Gln Val
 965 970 975

Leu Ala His Pro Val His Ala Pro Gly Ser Gly Gly Ala Ala Glu Gln
 980 985 990

Leu Ser Ser Lys Ser Arg Arg Arg Val Ser Ala Phe Pro Ser Ser Ile
 995 1000 1005

Pro Ala Glu Ser Leu Cys Pro Pro Leu Gln Gly Arg Arg Gln Gln
 1010 1015 1020

Lys Glu Gly Gln Glu Gly Ser His Ser Pro Val Pro Val Thr Arg
 1025 1030 1035

Leu Lys Asn Leu Ile Thr Cys Leu Asn Gly Ala Gly Glu Ile Asn
 1040 1045 1050

Asn Thr Thr Ser Leu Pro Glu Thr Val Arg Glu Trp Ser Leu Ser
 1055 1060 1065

Ser Gly Pro Ser Pro Leu Ala Gln Arg Arg Ser Val Gly Val Ile
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Pro Asn Ser Phe Leu Gln Thr Ser Ala Leu Ala Ser Ser Ile Gly
 1085 1090 1095

Arg Ser Phe His Leu Leu Arg Asn Gln Thr Arg Lys Ile Arg Cys
 1100 1105 1110

Asn Cys Ser His Gln Gly Arg Thr Leu Tyr Leu Val Cys Tyr Pro
 1115 1120 1125

Tyr Leu Leu Leu Thr Ser Leu Lys Gln Gln Gln Pro Thr Lys Arg
 1130 1135 1140

Cys Leu Glu Gln Ser Glu Leu Gln Val Leu Gln Ser Ser Ser Phe
 1145 1150 1155

Cys Pro Ala Thr Ser Ala Phe Lys Asn Gln Lys Lys Gly Gln Gly
 1160 1165 1170

Ala Gly Leu Leu Leu Thr Trp Ile Pro Lys Gln Gly Asp His Leu
 1175 1180 1185

Glu Leu Leu Gly Gln Arg Lys Glu Arg Thr Glu Pro Ala Ala Pro
 1190 1195 1200

Thr Pro Phe Ser His Met Pro Gln Ala Leu Ala Ala Leu Trp Thr
 1205 1210 1215

Gly Gly Gln Arg Ala His Glu Gln Leu Ala Arg Asp Gly Gln Pro
 1220 1225 1230

Asn Ser Thr Phe Pro Leu Leu Asp Gly Pro Gln His Leu Ser Asp
 1235 1240 1245

Leu Leu Ile Leu Gly Lys Gln Arg Leu Pro Ser Leu Ser Ile Ala
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Thr His Trp Trp Pro Ser Ser Thr Ser Glu Phe Leu Gln Pro Gly
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Arg Pro Leu Glu His
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<400> 5
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21

<210> 6
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120

cgttagcgcc ggatccccgc gggtagggcg gggcgggcgg cgccgtgggg atccccgggc	180
agccgagggc ccctgactcg gtcctcgcg gcgacatgga tcggatggcc agctccatga	240
agcaggtgcc caaccactg cccaaggtgc tgagccggcg cggggtcggc gctgggctgg	300
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ccattaatac gcaggaaagt ggctgtaaag gaaaaacatg ccag	404